RAW SEQUENCE LISTING

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Application Serial Number: 10 582,861
Source: If two
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PATENT APPLICATION: US/10/582,861   TIME: 11:13:21

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 3 <110> APPLICANT: Nobuo SAKAGUCHI et al.
 5 <120> TITLE OF INVENTION: ANTI-HIV ANTIBODY
 7 <130> FILE REFERENCE: 4456-0109PUS1
 9 <140> CURRENT APPLICATION NUMBER: US 10/582,861
10 <141> CURRENT FILING DATE: 2006-06-14
12 <150> PRIOR APPLICATION NUMBER: PCT/JP2004/003046
13 <151> PRIOR FILING DATE: 2004-03-09
15 <150> PRIOR APPLICATION NUMBER: JP 2003-418655
16 <151> PRIOR FILING DATE: 2003-12-16
18 <160> NUMBER OF SEQ ID NOS: 9
20 <170> SOFTWARE: PatentIn version 3.2
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 6429
24 <212> TYPE: DNA
25 <213> ORGANISM: Mus musclus
28 <220> FEATURE:
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30 <222> LOCATION: (384)..(6299)
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37 gtctttgccg agttgctgaa ggccttcact agtcttcgct cgaaggcgtc tgttaaccta    180
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43 cttcccctgg caacagataa tttaaagagg agagctactt gtgtatagtc catatttatt    360
45 gccttcagat aattggcttg aag atg cac ccg gtg aac ccc ttc gga ggc agc    413
46                          Met His Pro Val Asn Pro Phe Gly Gly Ser
47                           1               5                  10
49 agc cca agt gct ttt gcg gta tct tcc agc acc acg gga aca tat cag    461
50 Ser Pro Ser Ala Phe Ala Val Ser Ser Ser Thr Thr Gly Thr Tyr Gln
51                 15                  20                  25
53 act aaa tca cca ttt cga ttt ggc cag cct tcc ctt ttt gga cag aac    509
54 Thr Lys Ser Pro Phe Arg Phe Gly Gln Pro Ser Leu Phe Gly Gln Asn
55             30                  35                  40
57 agc aca ccc agc aag agc ctg gcg ttt tca caa gta cca agc ttt gca    557
58 Ser Thr Pro Ser Lys Ser Leu Ala Phe Ser Gln Val Pro Ser Phe Ala
59         45                  50                  55
61 aca ccc tct gga gga agc cat tct tcc tcc ttg cca gca ttt gga ctc    605
62 Thr Pro Ser Gly Gly Ser His Ser Ser Ser Leu Pro Ala Phe Gly Leu
63     60                  65                  70
65 acc caa acc tca agt gtg gga ctc ttc tct agt ctc gaa tcc aca cct    653
66 Thr Gln Thr Ser Ser Val Gly Leu Phe Ser Ser Leu Glu Ser Thr Pro
67 75                  80                  85                  90
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 73 agc ttt aag tca acc tct agc gtt ggg gtt ttc cca agt ggc gct act      749
 74 Ser Phe Lys Ser Thr Ser Ser Val Gly Val Phe Pro Ser Gly Ala Thr
 75             110                115                120
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 78 Phe Gly Pro Glu Thr Gly Glu Val Ala Gly Ser Gly Phe Arg Lys Thr
 79             125                130                135
 81 gaa ttc aag ttt aaa cct ctg gaa aat gca gtc ttc aaa ccg ata ccg      845
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 83         140                145                150
 85 ggg cct gag tca gag cca gaa aaa acc cag agc cag att tct tct gga      893
 86 Gly Pro Glu Ser Glu Pro Glu Lys Thr Gln Ser Gln Ile Ser Ser Gly
 87 155                160                165                170
 89 ttt ttt aca ttt tcc cat ccc gtt ggt agc ggg tct gga ggc ctg acc      941
 90 Phe Phe Thr Phe Ser His Pro Val Gly Ser Gly Ser Gly Gly Leu Thr
 91                 175                180                185
 93 cct ttt tct ttc cca cag gtg aca aat agt tcg gtg act agc tca agt      989
 94 Pro Phe Ser Phe Pro Gln Val Thr Asn Ser Ser Val Thr Ser Ser Ser
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 97 ttt atc ttt tcg aaa cca gtt act agt aat act cct gcc ttt gcc tct     1037
 98 Phe Ile Phe Ser Lys Pro Val Thr Ser Asn Thr Pro Ala Phe Ala Ser
 99         205                210                215
101 cct ttg tct aac caa aat gta gaa gaa gag aag agg gtt tct acg tca     1085
102 Pro Leu Ser Asn Gln Asn Val Glu Glu Glu Lys Arg Val Ser Thr Ser
103         220                225                230
105 gcg ttt gga agc tca aac agt agc ttc agt act ttc ccc aca gcg tca     1133
106 Ala Phe Gly Ser Ser Asn Ser Ser Phe Ser Thr Phe Pro Thr Ala Ser
107 235                240                245                250
109 cca gga tct ttg ggg gag ccc ttc cca gct aac aaa cca agc ctc cgc     1181
110 Pro Gly Ser Leu Gly Glu Pro Phe Pro Ala Asn Lys Pro Ser Leu Arg
111                 255                260                265
113 caa gga tgt gag gaa gcc atc tcc cag gtg gag cca ctt ccc acc ctc     1229
114 Gln Gly Cys Glu Glu Ala Ile Ser Gln Val Glu Pro Leu Pro Thr Leu
115             270                275                280
117 atg aag gga tta aag agg aaa gag gac cag gat cgc tcc ccg agg aga     1277
118 Met Lys Gly Leu Lys Arg Lys Glu Asp Gln Asp Arg Ser Pro Arg Arg
119         285                290                295
121 cat tgc cac gag gca gca gaa gac cct gat ccc ctg tcc agg ggc gac     1325
122 His Cys His Glu Ala Ala Glu Asp Pro Asp Pro Leu Ser Arg Gly Asp
123 300                305                310
125 cat ccc cca gat aaa cgg cca gtc cgc ctc aac aga ccc cgg gga ggt     1373
126 His Pro Pro Asp Lys Arg Pro Val Arg Leu Asn Arg Pro Arg Gly Gly
127 315                320                325                330
129 act ttg ttt ggc cgg aca ata cag gag gtc ttc aaa agc aat aaa gag     1421
130 Thr Leu Phe Gly Arg Thr Ile Gln Glu Val Phe Lys Ser Asn Lys Glu
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138 Pro Gly Glu Ser Asp His Ala Ala Val Pro Gly Gly Ser Gln Ser Thr
139         365             370             375
141 atg gta cct tcc cgc ctt cca gct gtg act aaa gag gaa gaa gaa agt      1565
142 Met Val Pro Ser Arg Leu Pro Ala Val Thr Lys Glu Glu Glu Glu Ser
143         380             385             390
145 aga gat gag aaa gaa gat tct ctc agg gga aag tct gtg cgc cag agt      1613
146 Arg Asp Glu Lys Glu Asp Ser Leu Arg Gly Lys Ser Val Arg Gln Ser
147 395             400             405             410
149 aag cga agg gaa gag tgg atc tac agc ctc ggg ggc gtg tct tct tta      1661
150 Lys Arg Arg Glu Glu Trp Ile Tyr Ser Leu Gly Gly Val Ser Ser Leu
151         415             420             425
153 gag ctc aca gcc atc cag tgc aag aac atc ccc gac tac ctc aac gac      1709
154 Glu Leu Thr Ala Ile Gln Cys Lys Asn Ile Pro Asp Tyr Leu Asn Asp
155         430             435             440
157 aga gcc atc ctg gag aaa cac ttc agc aaa atc gct aaa gtc cag cgg      1757
158 Arg Ala Ile Leu Glu Lys His Phe Ser Lys Ile Ala Lys Val Gln Arg
159         445             450             455
161 gtc ttc acc aga cgc agc aag aag ctc gcc gtg att cat ttt ttc gac      1805
162 Val Phe Thr Arg Arg Ser Lys Lys Leu Ala Val Ile His Phe Phe Asp
163         460             465             470
165 cac gca tcg gca gcc ctg gct agg aag aag ggg aaa ggt ctg cat aag      1853
166 His Ala Ser Ala Ala Leu Ala Arg Lys Lys Gly Lys Gly Leu His Lys
167 475             480             485             490
169 gac gtg gtt atc ttt tgg cac aag aag aaa ata agt ccc agc aag aaa      1901
170 Asp Val Val Ile Phe Trp His Lys Lys Lys Ile Ser Pro Ser Lys Lys
171         495             500             505
173 ctc ttt ccc ctg aag gag aag ctt ggt gag agt gaa gcc agc cag ggc      1949
174 Leu Phe Pro Leu Lys Glu Lys Leu Gly Glu Ser Glu Ala Ser Gln Gly
175         510             515             520
177 atc gag gac tcc ccc ttt cag cac tcg cct ctc agc aag ccc atc gtg      1997
178 Ile Glu Asp Ser Pro Phe Gln His Ser Pro Leu Ser Lys Pro Ile Val
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181 agg cct gca gcc ggc agc ctc ctc agc aaa agc tct cca gtg aag aag      2045
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183 540             545             550
185 ccg agt ctt ctg aag atg cac cag ttt gag gcg gat cct ttt gac tct      2093
186 Pro Ser Leu Leu Lys Met His Gln Phe Glu Ala Asp Pro Phe Asp Ser
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198 Leu Asp Gln Arg Asp Arg Ile Met Arg Gln Ala Arg Val Lys Arg Thr
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206 Pro Glu Lys Glu Arg Tyr Leu Arg Glu Thr Arg Ser Gln Leu Ser Val
207 635                   640                   645                   650
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210 Phe Glu Val Val Pro Gly Thr Asp Gln Val Asp His Ala Ala Ala Val
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213 aag gag tac agc cgg tcc tct gca gat cag gag gag ccc ctg cca cat    2429
214 Lys Glu Tyr Ser Arg Ser Ser Ala Asp Gln Glu Glu Pro Leu Pro His
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218 Glu Leu Arg Pro Ser Ala Val Leu Ser Arg Thr Met Asp Tyr Leu Val
219         685                   690                   695
221 acc cag atc atg gac caa aag gaa ggc agc ctt cgg gat tgg tat gac    2525
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227 715                   720                   725                   730
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230 His Leu Cys Asp Pro Leu Thr Val Ser Leu Ile Glu Lys Cys Thr Arg
231                   735                   740                   745
233 ttt cac att cac tgt gcc cac ttt atg tgt gag gag cct atg tct tcc    2669
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235                   750                   755                   760
237 ttt gat gcc aag atc aac aat gag aac atg acc aag tgt cta cag agt    2717
238 Phe Asp Ala Lys Ile Asn Asn Glu Asn Met Thr Lys Cys Leu Gln Ser
239         765                   770                   775
241 ctg aag gag atg tac cag gac ctg agg aac aag ggt gtt ttt tgt gcc    2765
242 Leu Lys Glu Met Tyr Gln Asp Leu Arg Asn Lys Gly Val Phe Cys Ala
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245 agt gaa gca gag ttt cag ggc tac aat gtc ctg ctt aat ctc aac aaa    2813
246 Ser Glu Ala Glu Phe Gln Gly Tyr Asn Val Leu Leu Asn Leu Asn Lys
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249 gga gac att ttg aga gaa gtg cag cag ttc cac cct gac gtt agg aac    2861
250 Gly Asp Ile Leu Arg Glu Val Gln Gln Phe His Pro Asp Val Arg Asn
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253 tcc cca gag gtg aac ttc gct gtc cag gct ttt gct gca ttg aac agc    2909
254 Ser Pro Glu Val Asn Phe Ala Val Gln Ala Phe Ala Ala Leu Asn Ser
255                   830                   835                   840
257 aat aat ttt gtg aga ttt ttc aaa ctg gtt cag tca gct tct tac ctg    2957
258 Asn Asn Phe Val Arg Phe Phe Lys Leu Val Gln Ser Ala Ser Tyr Leu
259         845                   850                   855
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263     860                   865                   870
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265 ctc cgg gca ctc aat gtt gct tat act gta agc aca cag cgc tct acc       3053
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269 gtc ttc ccc ctg gat ggt gtc gtc cgc atg ctg ctg ttc aga gat agt       3101
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271             895             900             905
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278 Gly Cys Val Glu Leu Asn Arg Ser Ala Phe Leu Glu Pro Glu Gly Leu
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293 ctg gct acg gag ctg ccc atc agc act cag aga gct ggt gga gac cca       3389
294 Leu Ala Thr Glu Leu Pro Ile Ser Thr Gln Arg Ala Gly Gly Asp Pro
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319         1080             1085             1090
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322 Ser Ala Gly Ala Ala Tyr Val Ala Ala Ala Leu Gly Val Ser Asn
323         1095             1100             1105
325 gct gct gtg gag gat ctg att act gct gcg acc acg ggc att ctg       3749
326 Ala Ala Val Glu Asp Leu Ile Thr Ala Ala Thr Thr Gly Ile Leu
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